SEQUENCE LISTING

<110> University of Rochester Fay, Philip J. Wakabayashi, Hironao <120> RECOMBINANT FACTOR VIII HAVING INCREASED SPECIFIC ACTIVITY <130> 176/61701 <140> <141> <150> 60/526,664 <151> 2003-12-03 <160> 7 <170> PatentIn Ver. 2.1 <210> 1 <211> 6999 <212> DNA <213> Human <400> 1 gccaccagaa gatactacct gggtgcagtg gaactgtcat gggactatat gcaaagtgat 60 ctcggtgagc tgcctgtgga cgcaagattt cctcctagag tgccaaaatc ttttccattc 120 aacacctcag tcgtgtacaa aaagactctg tttgtagaat tcacggatca ccttttcaac 180 atcgctaagc caaggccacc ctggatgggt ctgctaggtc ctaccatcca ggctgaggtt 240 tatgatacag tggtcattac acttaagaac atggcttccc atcctgtcag tcttcatgct 300 gttggtgtat cctactggaa agcttctgag ggagctgaat atgatgatca gaccagtcaa 360 agggagaaag aagatgataa agtcttccct ggtggaagcc atacatatgt ctggcaggtc 420 ctgaaagaga atggtccaat ggcctctgac ccactgtgcc ttacctactc atatctttct 480 catgtggacc tggtaaaaga cttgaattca ggcctcattg gagccctact agtatgtaga 540 gaagggagtc tggccaagga aaagacacag accttgcaca aatttatact actttttgct 600 gtatttgatg aagggaaaag ttggcactca gaaacaaaga actccttgat gcaggatagg 660 gatgctgcat ctgctcgggc ctggcctaaa atgcacacag tcaatggtta tgtaaacagg 720 tctctgccag gtctgattgg atgccacagg aaatcagtct attggcatgt gattggaatg 780 ggcaccactc ctgaagtgca ctcaatattc ctcgaaggtc acacatttct tgtgaggaac 840 catcgccagg cgtccttgga aatctcgcca ataactttcc ttactgctca aacactcttg 900 atggaccttg gacagtttct actgttttgt catatctctt cccaccaaca tgatggcatg 960 gaagettatg teaaagtaga eagetgteea gaggaaceee aactaegaat gaaaaataat 1020 gaagaagcgg aagactatga tgatgatctt actgattctg aaatggatgt ggtcaggttt 1080 gatgatgaca actotoctto otttatocaa attogotoag ttgocaagaa goatootaaa 1140 acttgggtac attacattgc tgctgaagag gaggactggg actatgctcc cttagtcctc 1200

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Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val 65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val 85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala 100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val 115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn 130 135 140

Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser 145 150 155 160

His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu 165 170 175

Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu 180 185 190

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His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser

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Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile

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- Val Ser Ser Ser Asp Leu Leu Met Leu Leu Arg Gln Ser Pro Thr Pro 785 790 795 800
- His Gly Leu Ser Leu Ser Asp Leu Gln Glu Ala Lys Tyr Glu Thr Phe 805 810 815
- Ser Asp Asp Pro Ser Pro Gly Ala Ile Asp Ser Asn Asn Ser Leu Ser 820 825 830
- Glu Met Thr His Phe Arg Pro Gln Leu His His Ser Gly Asp Met Val 835 840 845
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- Tyr Asp Ser Gln Leu Asp Thr Thr Leu Phe Gly Lys Lys Ser Ser Pro 915 920 925
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- Ser Lys Leu Leu Glu Ser Gly Leu Met Asn Ser Gln Glu Ser Ser Trp 945 950 955 960
- Gly Lys Asn Val Ser Ser Thr Glu Ser Gly Arg Leu Phe Lys Gly Lys 965 970 975
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- Val Cys His Thr Asn Thr Leu Asn Pro Ala His Gly Arg Gln Val Thr 1860 1865 1870
- Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe Asp Glu Thr Lys Ser 1875 1880 1885
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2005 2010 2015

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10

15

Asp